

SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT: KIKUCHI, KOKICHI
SATO, NORIYUKI
SAHARA, HIROMITSU
YASOJIMA, TAKAHIRO
WADA, YOSHIMASA
SUZUKI, MANABU
HAMURO, JUNJI

(ii) TITLE OF INVENTION: PEPTIDE CAPABLE OF INDUCING IMMUNE
RESPONSE TO HUMAN GASTRIC CANCER AND AGENT FOR PREVENTING
OR TREATING HUMAN GASTRIC CANCER, CONTAINING THE PEPTIDE

(iii) NUMBER OF SEQUENCES: 4

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: OBLON, SPIVAK, McCLELLAND, MAIER & NEUSTADT,
P.C.
(B) STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
(C) CITY: ARLINGTON
(D) STATE: VA
(E) COUNTRY: USA
(F) ZIP: 22202

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.30

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: US 08/723116
(B) FILING DATE: 30-SEP-1996
(C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: JP 253491/1995
(B) FILING DATE: 29-SEP-1995

(viii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: JP 217140/1996
(B) FILING DATE: 19-AUG-1996

(ix) ATTORNEY/AGENT INFORMATION:

(A) NAME: OBLON, NORMAN F.
(B) REGISTRATION NUMBER: 24,618
(C) REFERENCE/DOCKET NUMBER: 10-821-0X

(x) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: 703-413-3000
(B) TELEFAX: 703-413-2220

(2) INFORMATION FOR SEQ ID NO:1:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 10 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(vi) ORIGINAL SOURCE:

(A) ORGANISM: HUMAN

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

Tyr Ser Trp Met Asp Ile Ser Cys Trp Ile
1 5 10

(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 9 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(vi) ORIGINAL SOURCE:

(A) ORGANISM: HUMAN

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Tyr Ser Trp Met Asp Ile Ser Cys Trp
1 5 10

(2) INFORMATION FOR SEQ ID NO:3:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 8 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(vi) ORIGINAL SOURCE:

(A) ORGANISM: HUMAN

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

Tyr Ser Trp Met Asp Ile Ser Cys
1 5

(2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 7 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(vi) ORIGINAL SOURCE:

(A) ORGANISM: HUMAN

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

Tyr Ser Trp Met Asp Ile Ser
1 5